NEW SHEET

1	MAPVAVWAAL	AVGLELWAAA	<u>HA</u> LPAQVAFT	PYAPEPGSTC	RLREYYDQTA
51	QMCCSKCSPG	QHAKVFCTKT	SDTVCDSCED	STYTQLWNWV	PECLSCGSRC
101	SSDQVETQAC	TREQURICTO	RPGWYCALSK	QEGCRLCAPL	RKCRPGFGVA
151	RPGTETSDVV	CKPCAPGTFS	NTTSSTDICR	PHQICNVVAI	PG <u>NAS</u> HDAVC
201	TSTSPTRSMA	PGAVHLPQPV	STRSQHTQPT	PEPSTAPSTS	FLLPMGPSPP
251	AEGSTGDFAL	PVGLIVGVTA	LGLLIIGVVN	CVIMTQVKKK	PLCLQREAKV
301	PHLPADKARG	TQGPEQQHLL	ITAPSSSSSS	LESSASALDR	RAPTRNQPQA
351	PGVEASGAGE	ARASTGSSDS	SPGGHGTQV <u>N</u>	VTCIVNVCSS	SDHSSQCSSQ
401	ASSTMGDTDS	SPSESPKDEQ	VPFSKEECAF	RSQLETPETL	LGSTEEKPLP
451	LGVPDAGMKP	5			

FIGURE 5